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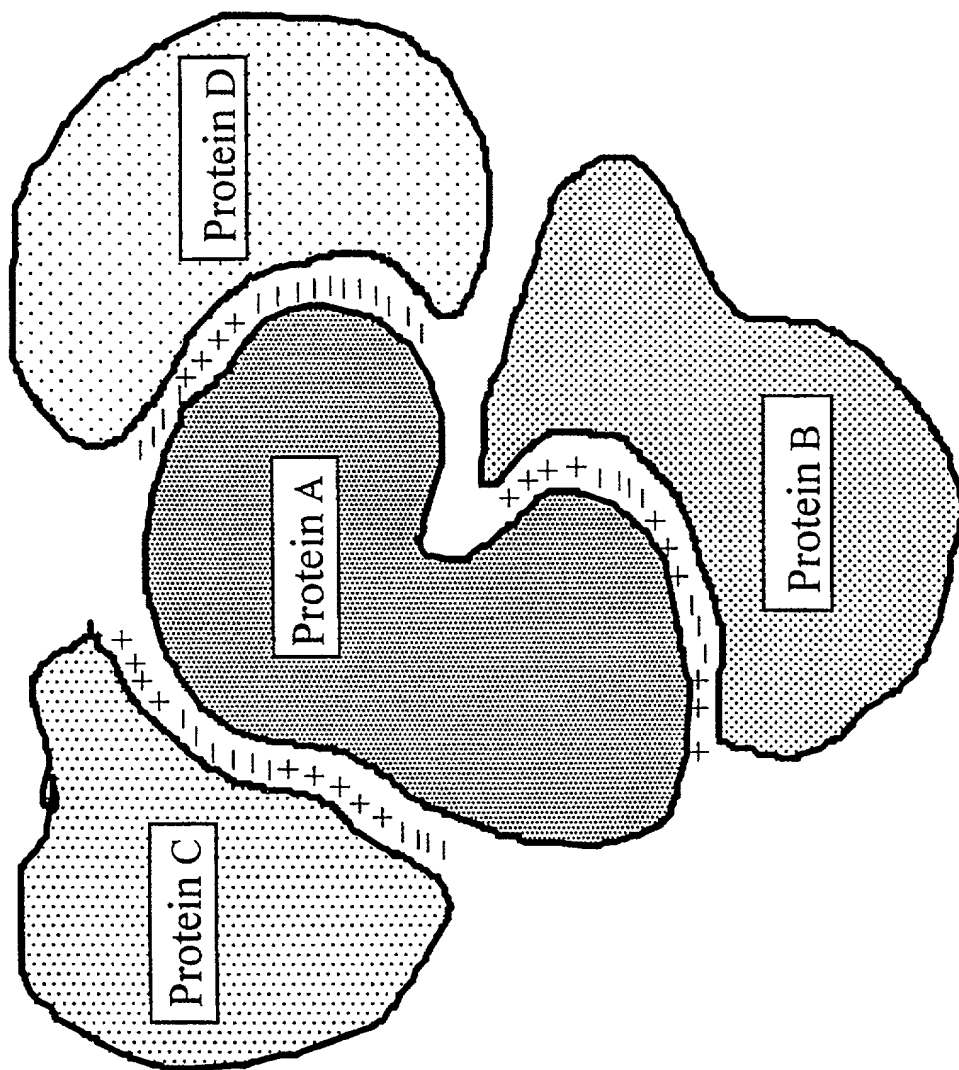
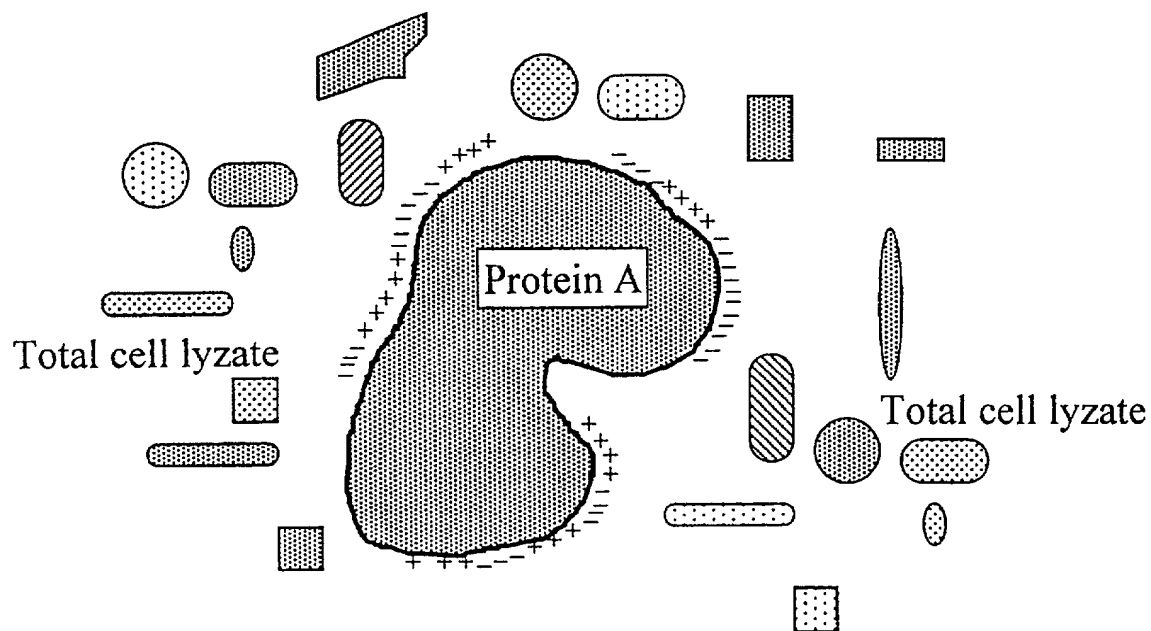
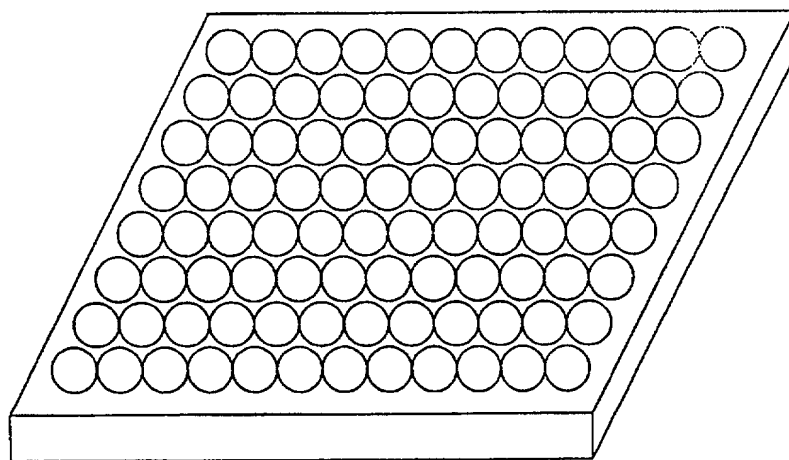


FIG. 1

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FIG. 2AFIG. 2B

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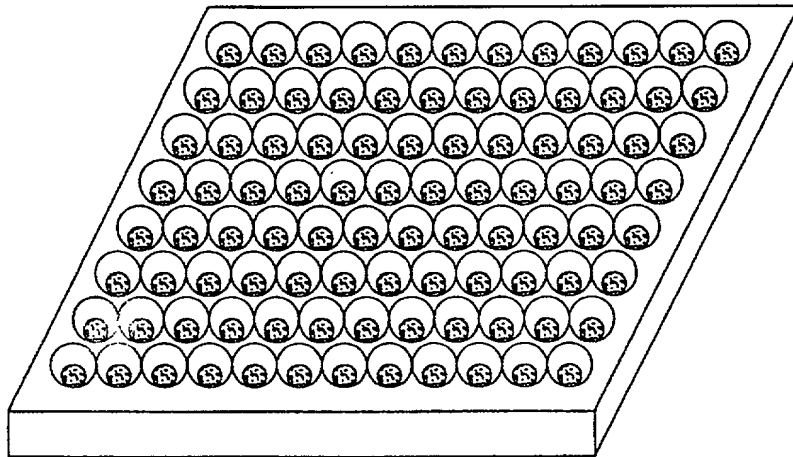


FIG. 2C

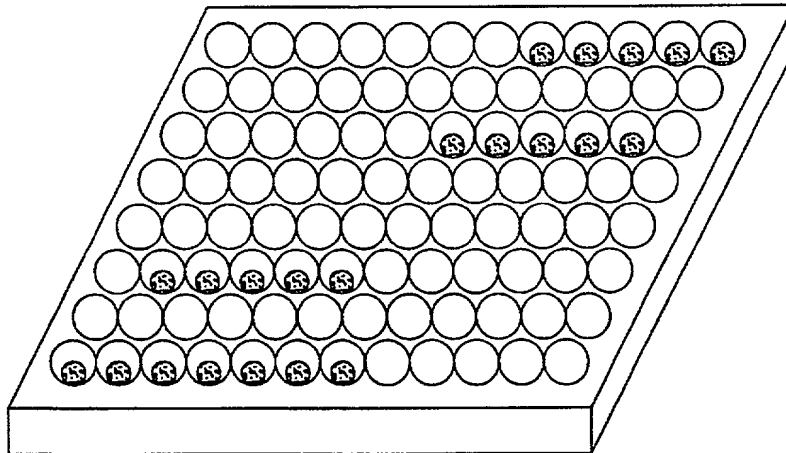


FIG. 2D

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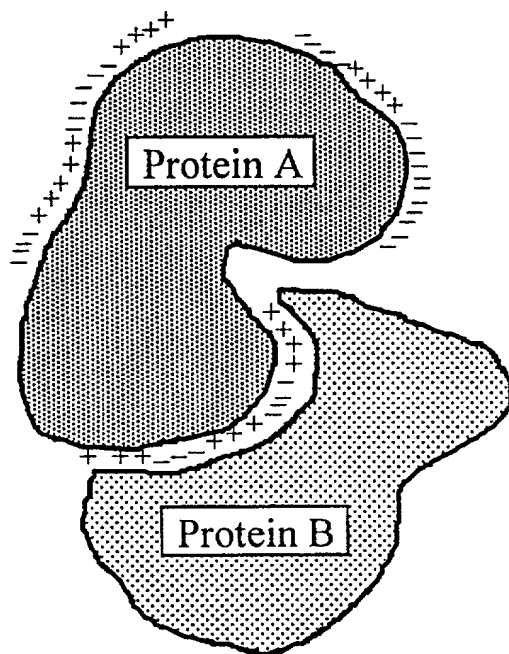


FIG. 3A

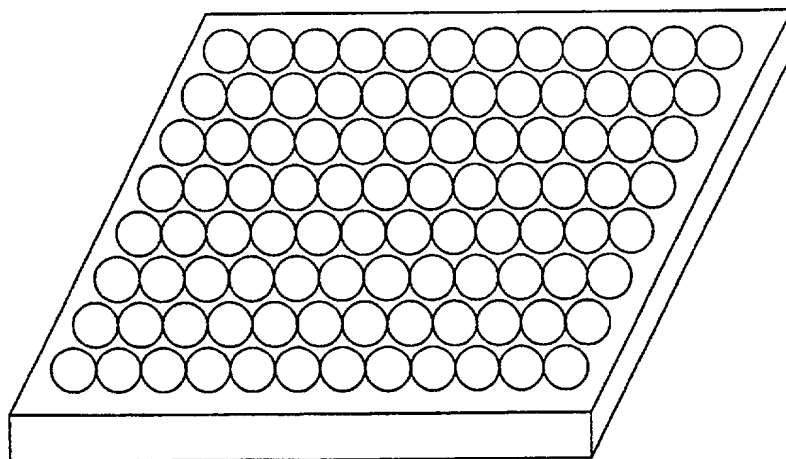


FIG. 3B

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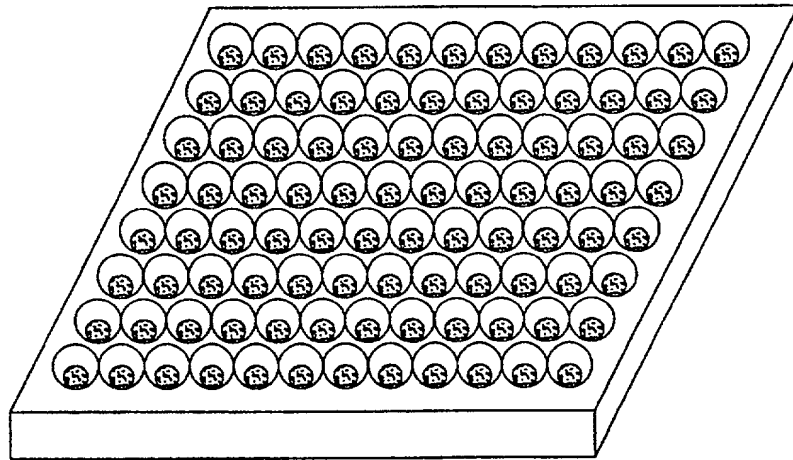


FIG. 3C

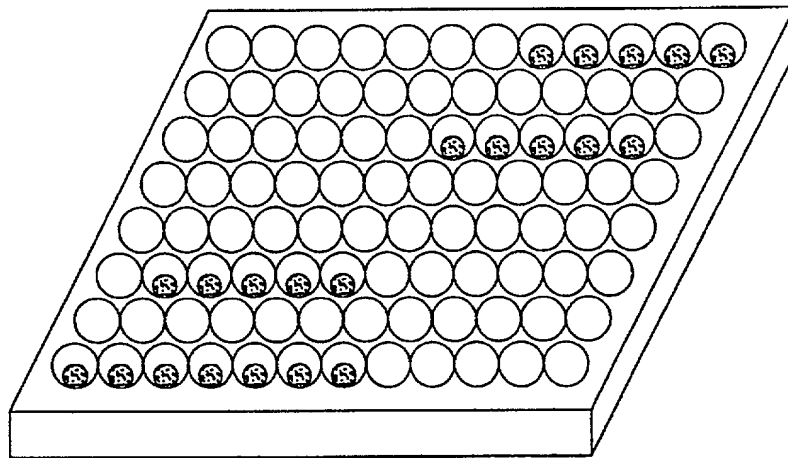


FIG. 3D

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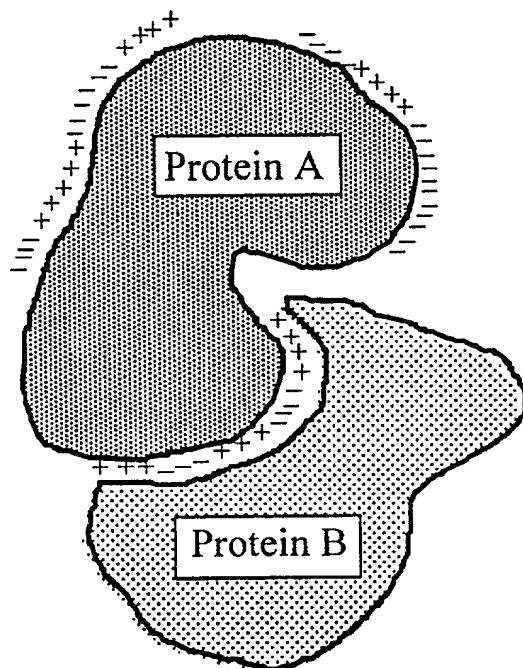


FIG. 4A

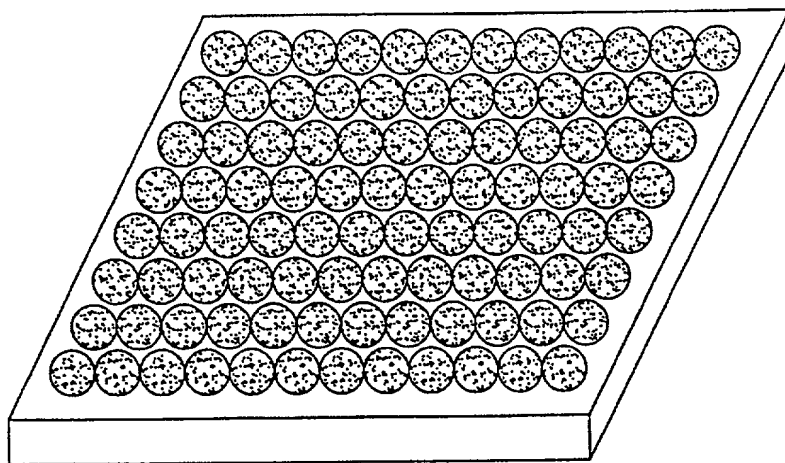
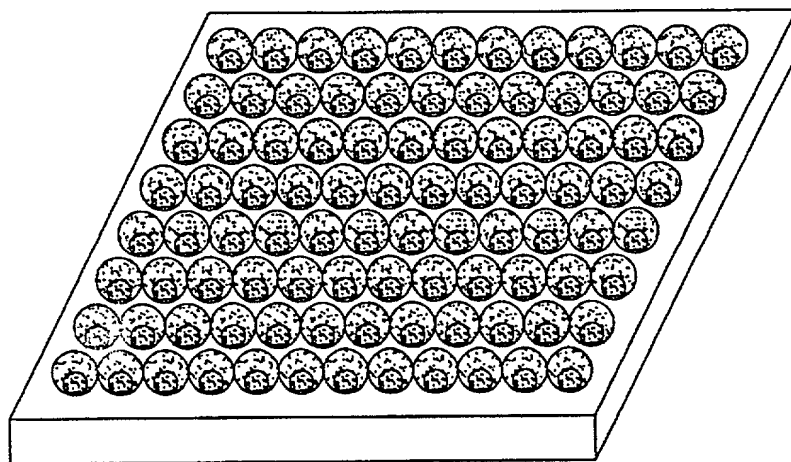
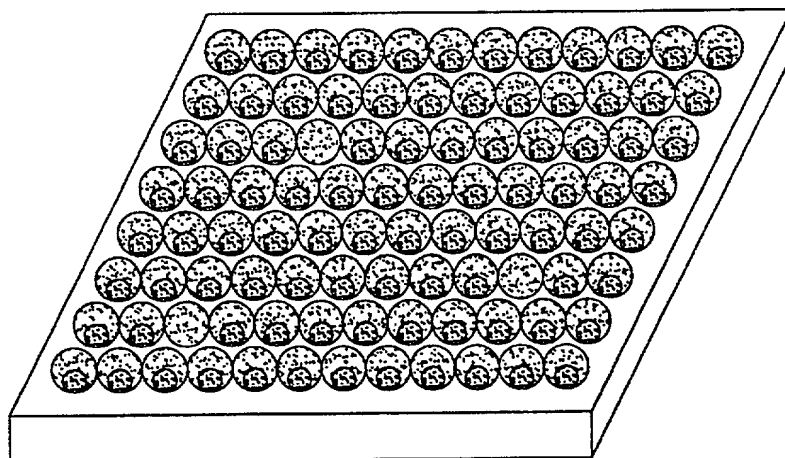


FIG. 4B

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FE - 4C



FE - 4D

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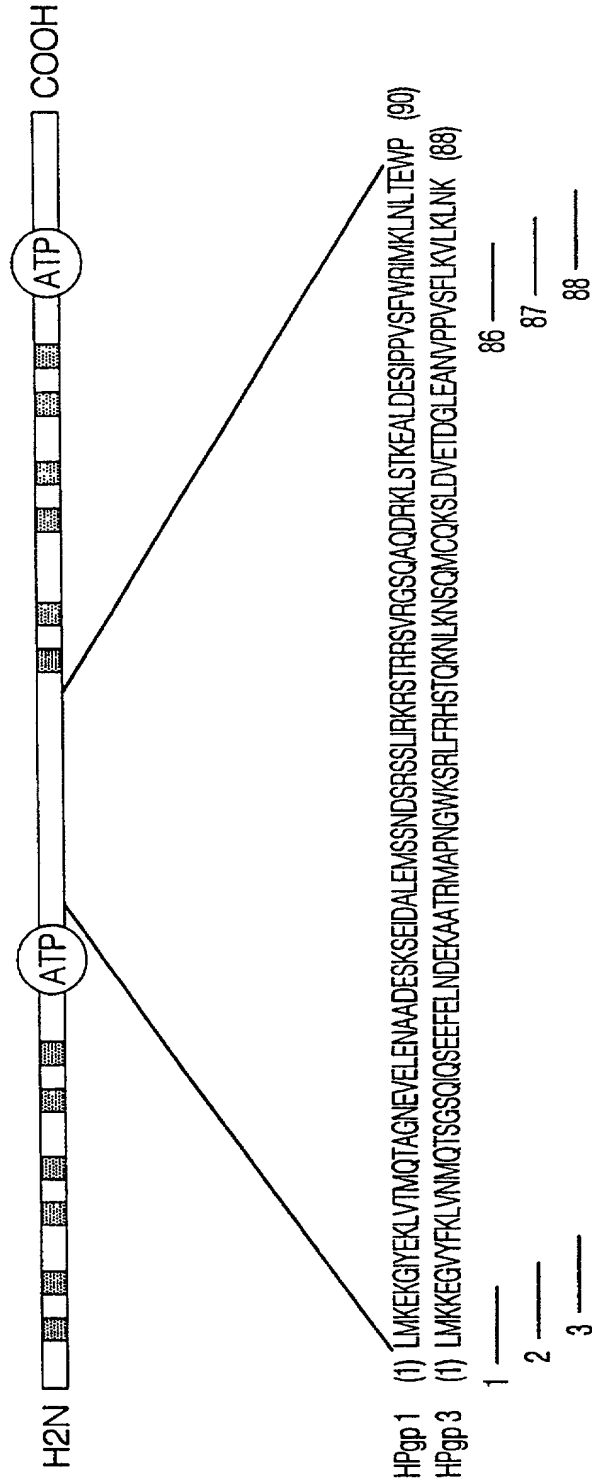


FIG. 5

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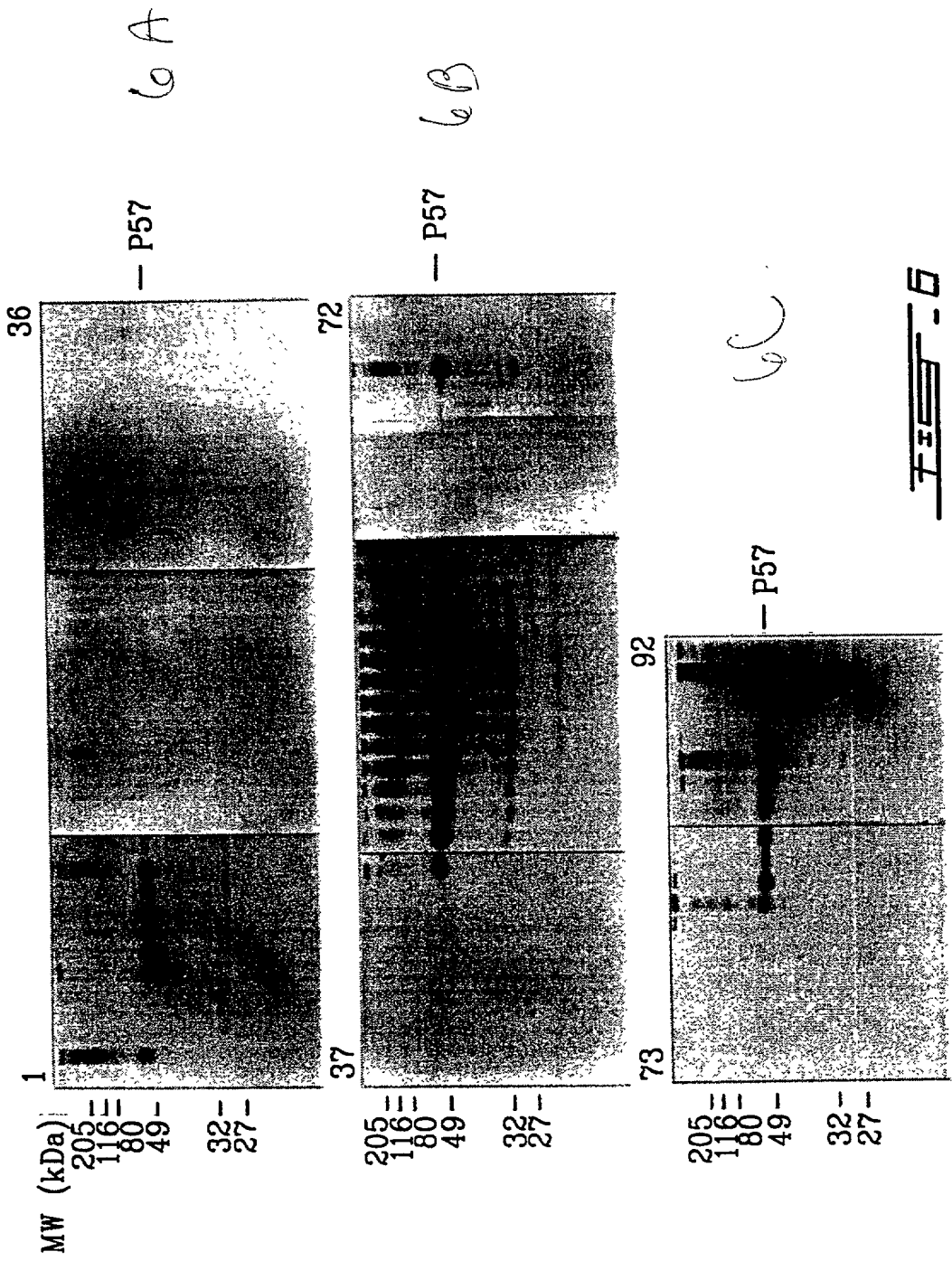


Figure 6

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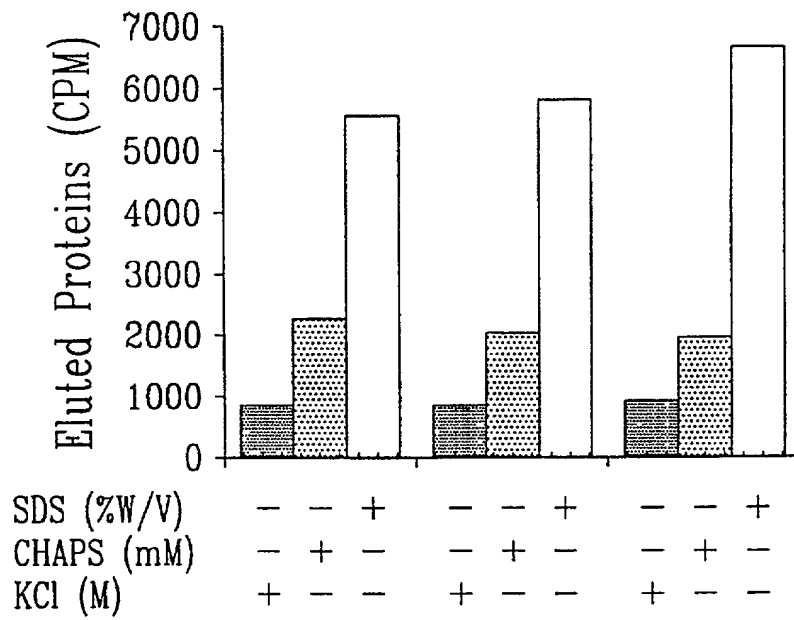
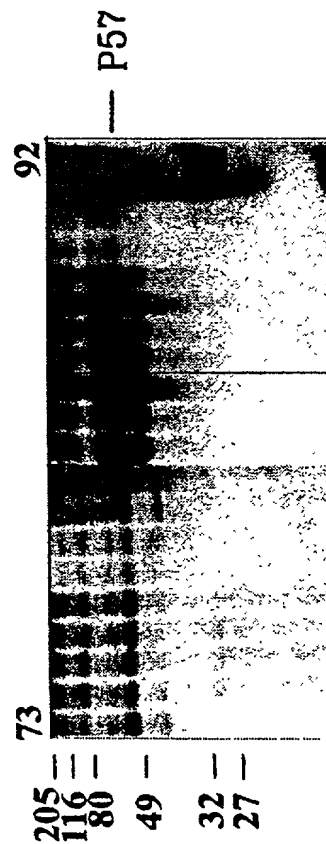
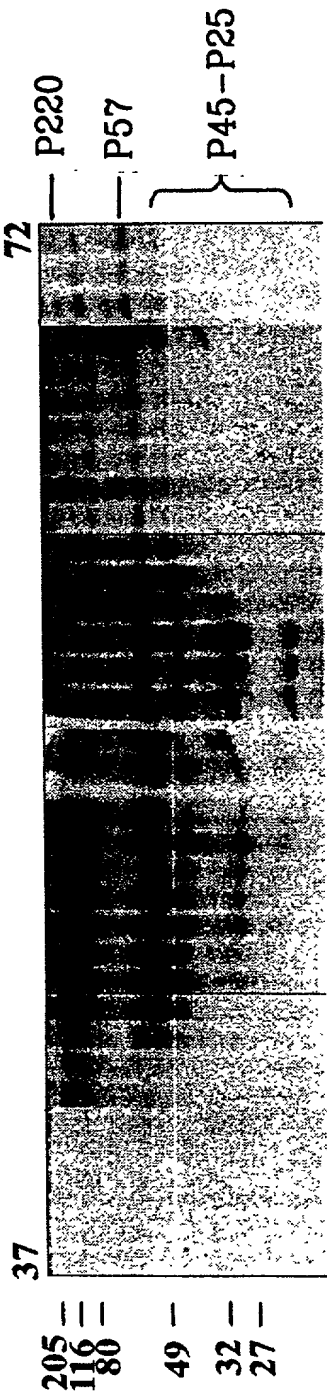
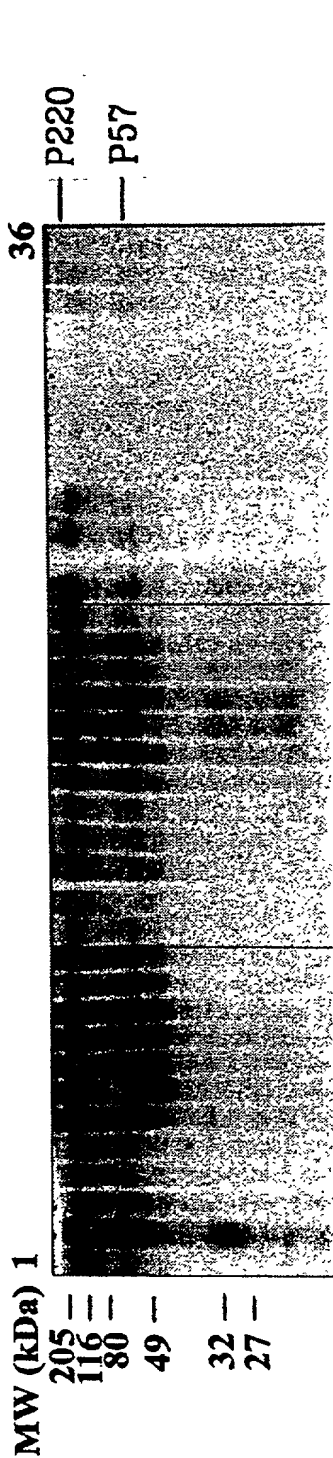


Fig. 7

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FILE - B

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FIG. 1. 01000000

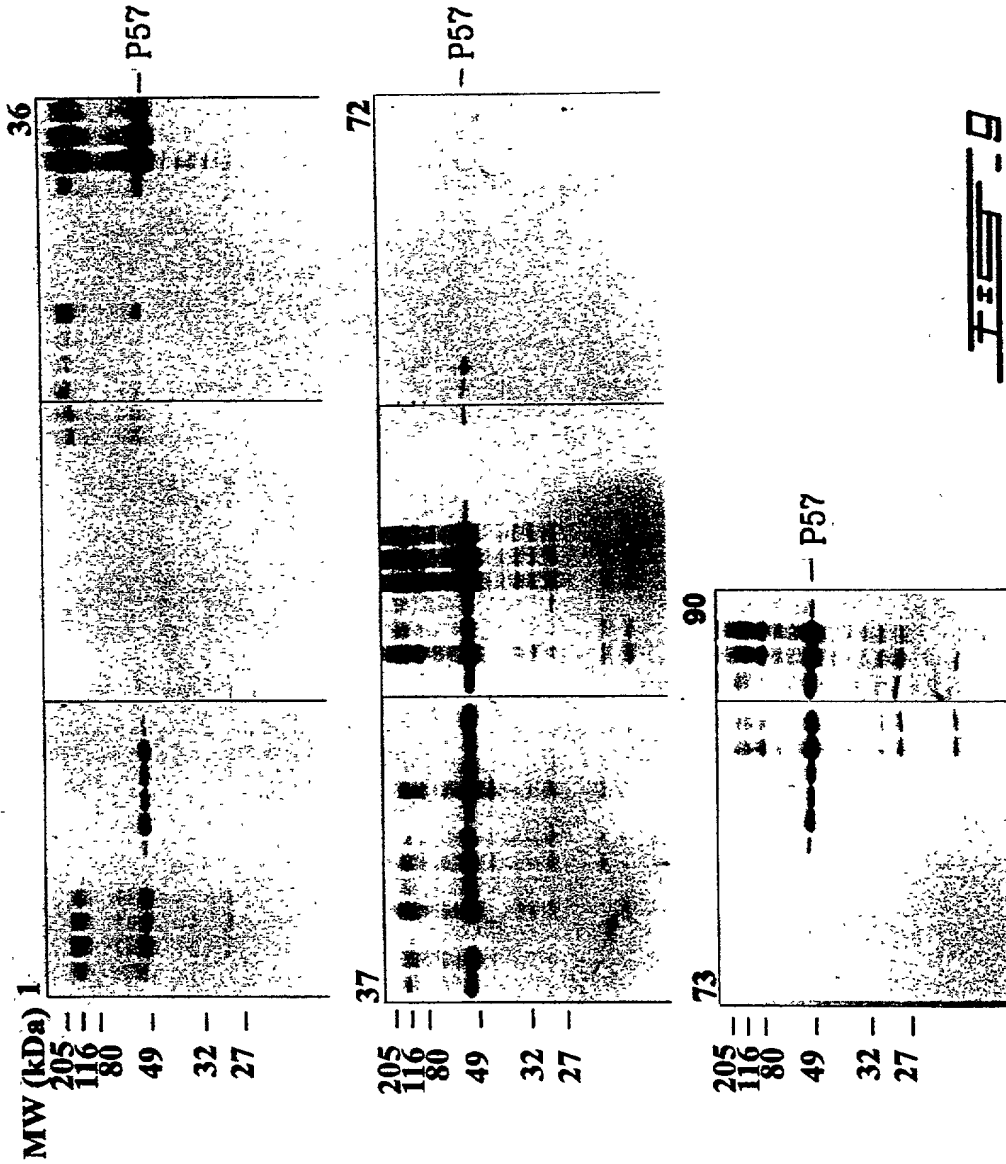


FIG. 1. 01000000

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Alignment Of Amino Acid Sequences Of Human P-gp3 And P-gp1 Linker Domains

618 **LMKKEGVYFKLVNMQTSGSQIQSEEF--ELNDEKAAATRMAPNGWKSRLFR-HSTQKNLKN SQM** 677 P-gp3
 LMK++G+YFKLV MQT+G++++ E E A M+ N +S L R ST+++++

615 **LMKEKGIYFKLVTMTAGNEVELENAADESKSEIDALEMSSNDSRSSLI RKRSTRRSV RGSQA** 677 P-gp1

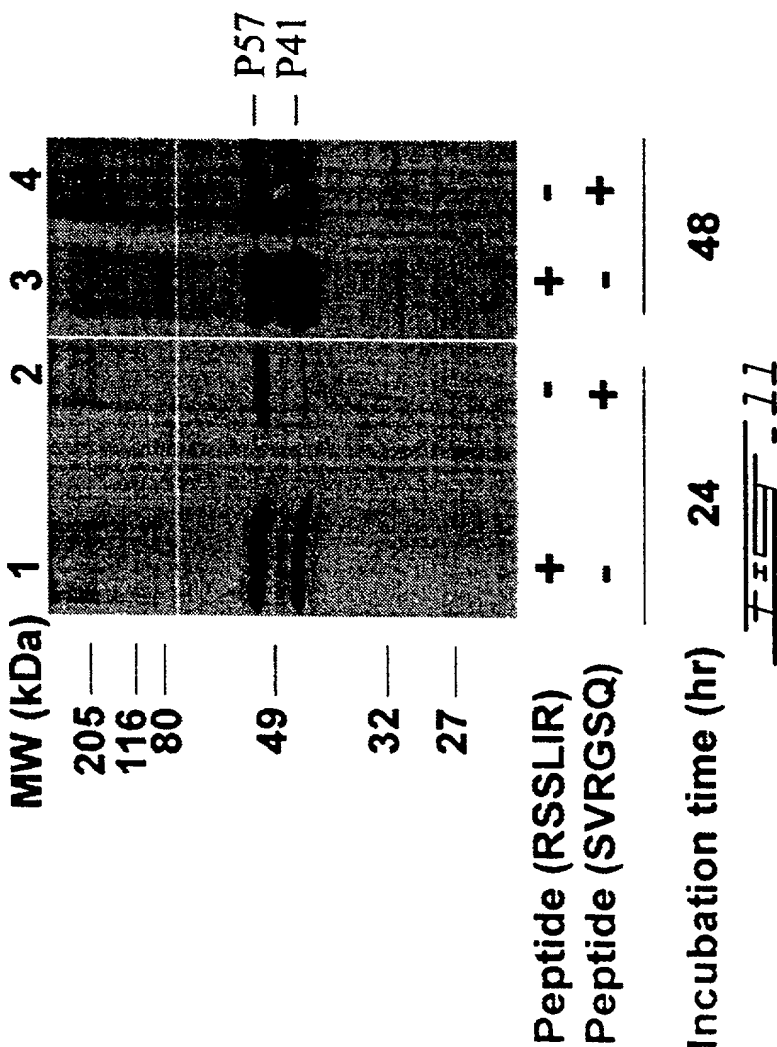
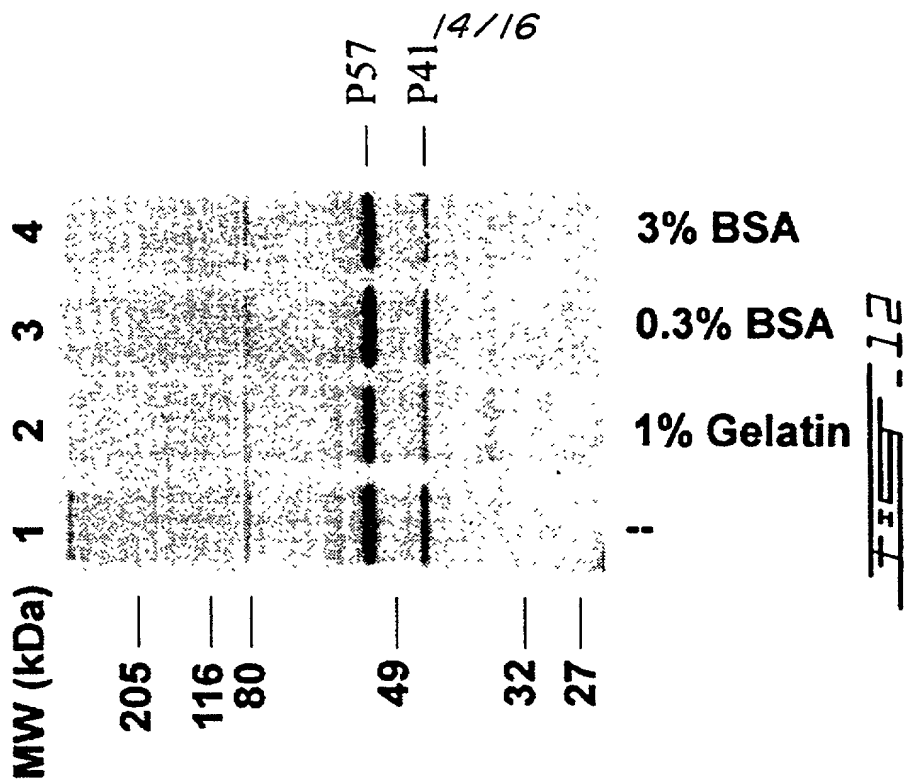
678 **CQKSLDVE TDGLEANVPVVSFLKVLKLNKTEWP** 710 P-gp3

+ L + + L+ ++PPVSF +++KLN TEWP

678 **QDRKLSTK-EALDESIPPVSFWRIMKLNLT EWP** 709 P-gp1

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FIG. 10



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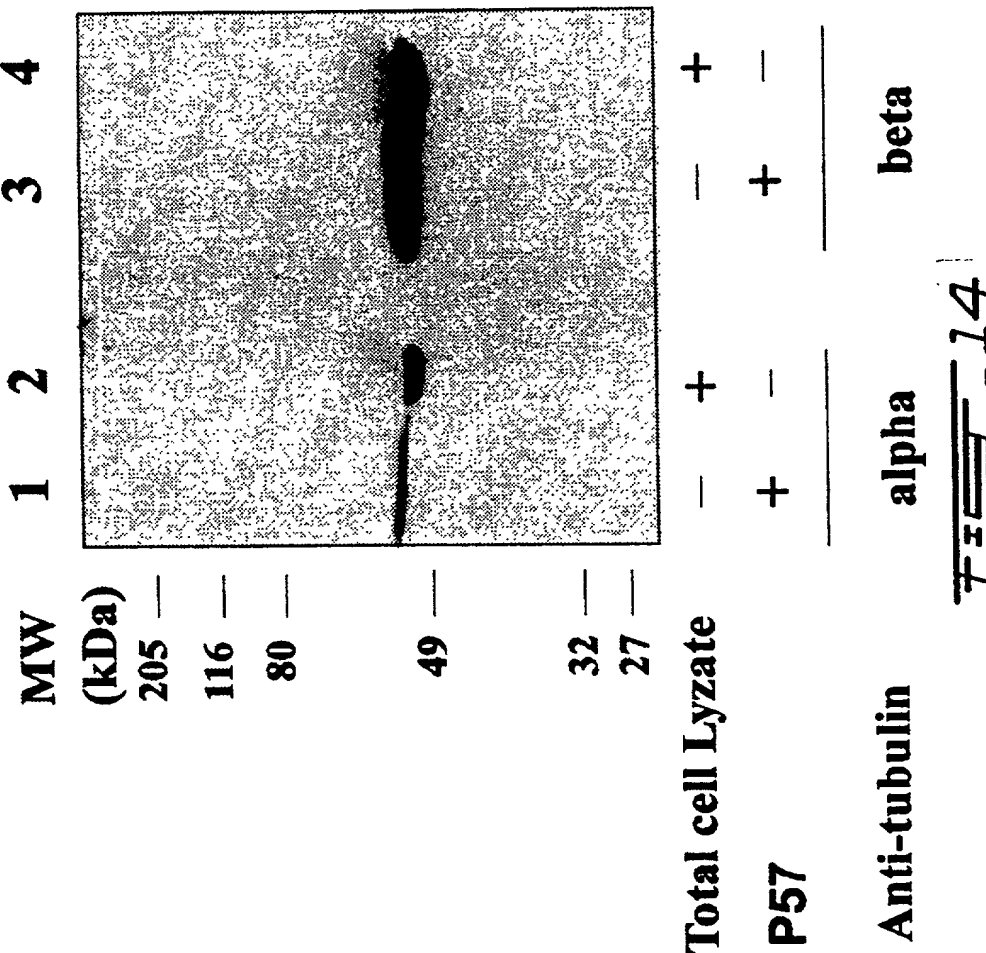


Fig. 14

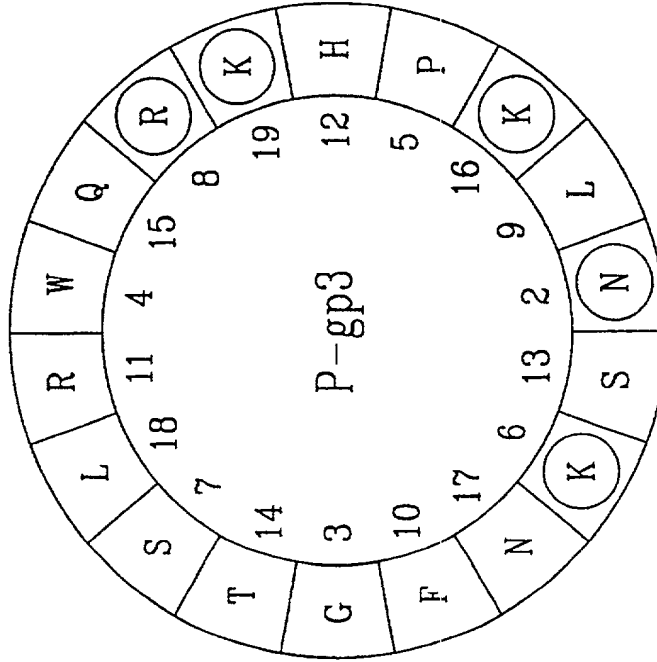


Fig. 15

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Helical Wheel Presentation

NGWKSRLFRHSTQKNLK



SRSSLIRKRSTRRSVRGS

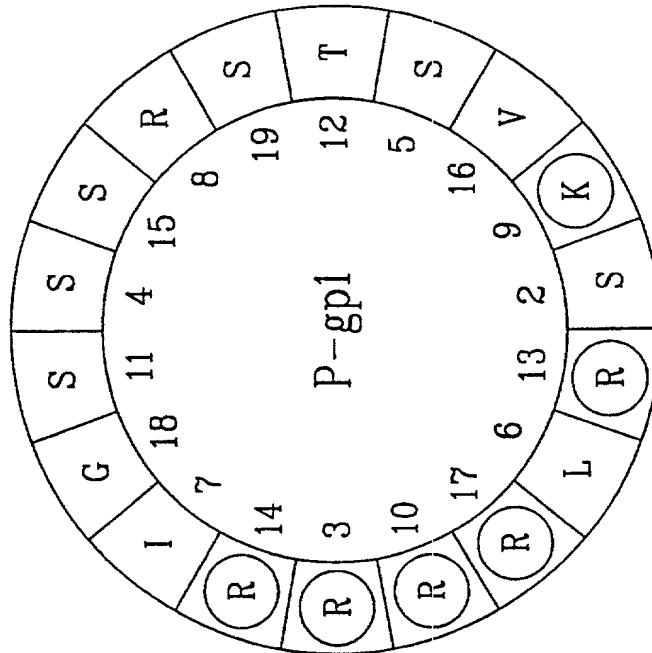


Fig. 15